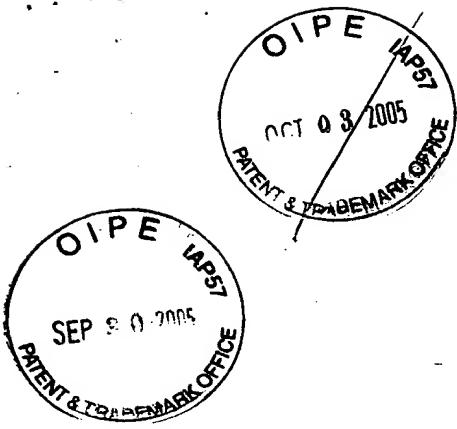


5/10



(SEQ ID NO: 37) AIR-1: 299 CAVCORDGGELICCDGCPRAFHLLACILSPPRLREIPSGTWRCSSC 340  
(SEQ ID NO: 38) AIR-1: 434 .G..G..TDVLR.TH.AA..WR.HF.AGTSR.GTGL..R.. 475  
(SEQ ID NO: 39) Mi-2 : 373 .E..QQ..I.L..T...Y.MV..D.DMEKA.E.K.S.PH. 414  
(SEQ ID NO: 40) Mi-2 : 452 .R..K.....T..SSY.IH..N..P...N.E.L.PR. 493  
(SEQ ID NO: 41) TIF1 : 791 ...QN.....EK..KV...S.HV.T.TNF...E.I.TF. 832  
consensus C C H C C C C

FIG. 5